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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/780,224A

DATE: 06/07/2001
TIME: 13:54:10

Input Set : **A:\208859.txt**
Output Set: **C:\CRF3\06072001\I780224A.raw**

ENTERED

```
 3 <110> APPLICANT: Roelvink, Petrus W
 4         Kovesdi, Imre
 5         Wickham, Thomas J
 7 <120> TITLE OF INVENTION: ADENOVIRAL CAPSID CONTAINING CHIMERIC PROTEIN IX
 9 <130> FILE REFERENCE: 208859
11 <140> CURRENT APPLICATION NUMBER: US 09/780,224A
12 <141> CURRENT FILING DATE: 2001-02-09
14 <150> PRIOR APPLICATION NUMBER: US 60/181,163
15 <151> PRIOR FILING DATE: 2000-02-09
17 <160> NUMBER OF SEQ ID NOS: 13
19 <170> SOFTWARE: PatentIn Ver. 2.1
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 144
23 <212> TYPE: PRT
24 <213> ORGANISM: Adenovirus
26 <400> SEQUENCE: 1
27 Met Asn Gly Thr Thr Gln Asn Asn Ala Ala Leu Phe Asp Gly Gly Val
28   1               5                   10                  15
30 Phe Ser Pro Tyr Leu Thr Ser Arg Leu Pro Tyr Trp Ala Gly Val Arg
31              20                  25                  30
33 Gln Asn Val Val Gly Ser Thr Val Asp Gly Arg Pro Val Ala Pro Ala
34           35                  40                  45
36 Asn Ser Ser Thr Leu Thr Tyr Ala Thr Ile Gly Pro Ser Pro Leu Asp
37        50                  55                  60
39 Thr Ala Ala Ala Ala Ala Ala Ser Ala Ala Ala Ser Thr Ala Arg Ser
40  65                  70                  75                  80
42 Met Ala Ala Asp Phe Ser Phe Tyr Asn His Leu Ala Ser Asn Ala Val
43              85                  90                  95
45 Thr Arg Thr Ala Val Arg Glu Asp Ile Leu Thr Val Met Leu Ala Lys
46             100                 105                 110
48 Leu Glu Thr Leu Thr Ala Gln Leu Glu Glu Leu Ser Gln Lys Val Glu
49          115                 120                 125
51 Glu Leu Ala Asp Ala Thr Thr His Thr Pro Ala Gln Pro Val Thr Gln
52       130                 135                 140
54 <210> SEQ ID NO: 2
55 <211> LENGTH: 125
56 <212> TYPE: PRT
57 <213> ORGANISM: Adenovirus
59 <400> SEQUENCE: 2
60 Met Ala Glu Glu Gly Arg Ile Tyr Val Pro Tyr Val Thr Ala Arg Leu
61   1               5                   10                  15
63 Pro Lys Trp Ser Gly Ser Val Gln Asp Lys Thr Gly Ser Asn Met Leu
64              20                  25                  30
66 Gly Gly Val Val Leu Pro Pro Asn Ser Gln Ala His Arg Thr Glu Thr
67           35                  40                  45
69 Val Gly Thr Glu Ala Thr Arg Asp Asn Leu His Ala Glu Gly Ala Arg
70        50                  55                  60
```

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72 Arg Pro Glu Asp Gln Thr Pro Tyr Met Ile Leu Val Glu Asp Ser Leu
73  65                  70                  75                  80
75 Gly Gly Leu Lys Arg Arg Met Asp Leu Leu Glu Glu Ser Asn Gln Gln
76                  85                  90                  95
78 Leu Leu Ala Thr Leu Asn Arg Leu Arg Thr Gly Leu Ala Ala Tyr Val
79                 100                 105                 110
81 Gln Ala Asn Leu Val Gly Gly Gln Val Asn Pro Phe Val
82                 115                 120                 125
84 <210> SEQ ID NO: 3
85 <211> LENGTH: 125
86 <212> TYPE: PRT
87 <213> ORGANISM: Adenovirus
89 <400> SEQUENCE: 3
90 Met Ala Glu Glu Gly Arg Ile Tyr Val Pro Tyr Val Thr Ala Arg Leu
91  1                   5                  10                  15
93 Pro Lys Trp Ser Gly Ser Val Gln Asp Lys Thr Gly Ser Asn Met Leu
94                  20                  25                  30
96 Gly Gly Val Val Leu Pro Pro Asn Ser Gln Ala His Arg Thr Glu Thr
97                  35                  40                  45
99 Val Gly Thr Glu Ala Thr Arg Asp Asn Leu His Ala Glu Gly Ala Arg
100                 50                  55                  60
102 Arg Pro Glu Asp Gln Thr Pro Tyr Met Ile Leu Val Glu Asp Ser Leu
103  65                 70                  75                  80
105 Gly Gly Leu Lys Arg Arg Met Asp Leu Leu Glu Glu Ser Asn Gln Gln
106                 85                  90                  95
108 Leu Leu Ala Thr Leu Asn Arg Leu Arg Thr Gly Leu Ala Ala Tyr Val
109                100                 105                 110
111 Gln Ala Asn Leu Val Gly Gly Gln Val Asn Pro Phe Val
112                115                 120                 125
114 <210> SEQ ID NO: 4
115 <211> LENGTH: 140
116 <212> TYPE: PRT
117 <213> ORGANISM: Adenovirus
119 <400> SEQUENCE: 4
120 Met Ser Ala Asn Ser Phe Asp Gly Ser Ile Val Ser Ser Tyr Leu Thr
121  1                   5                  10                  15
123 Thr Arg Met Pro Pro Trp Ala Gly Val Arg Gln Asn Val Met Gly Ser
124                 20                  25                  30
126 Ser Ile Asp Gly Arg Pro Val Leu Pro Ala Asn Ser Thr Thr Leu Thr
127                 35                  40                  45
129 Tyr Glu Thr Val Ser Gly Thr Pro Leu Glu Thr Ala Ala Ser Ala Ala
130                 50                  55                  60
132 Ala Ser Ala Ala Ala Ala Thr Ala Arg Gly Ile Val Thr Asp Phe Ala
133  65                 70                  75                  80
135 Phe Leu Ser Pro Leu Ala Ser Ser Ala Ala Ser Arg Ser Ser Ala Arg
136                 85                  90                  95
138 Asp Asp Lys Leu Thr Ala Leu Leu Ala Gln Leu Asp Ser Leu Thr Arg
139                100                 105                 110
141 Glu Leu Asn Val Val Ser Gln Gln Leu Leu Asp Leu Arg Gln Gln Val
```

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142          115                   120                    125
144 Ser Ala Leu Lys Ala Ser Ser Pro Pro Asn Ala Val
145      130             135             140
147 <210> SEQ ID NO: 5
148 <211> LENGTH: 140
149 <212> TYPE: PRT
150 <213> ORGANISM: Adenovirus
152 <400> SEQUENCE: 5
153 Met Ser Thr Asn Ser Phe Asp Gly Ser Ile Val Ser Ser Tyr Leu Thr
154   1                   5                   10                  15
156 Thr Arg Met Pro Pro Trp Ala Gly Val Arg Gln Asn Val Met Gly Ser
157             20                  25                  30
159 Ser Ile Asp Gly Arg Pro Val Leu Pro Ala Asn Ser Thr Thr Leu Thr
160             35                  40                  45
162 Tyr Glu Thr Val Ser Gly Thr Pro Leu Glu Thr Ala Ala Ser Ala Ala
163      50                  55                  60
165 Ala Ser Ala Ala Ala Ala Thr Ala Arg Gly Ile Val Thr Asp Phe Ala
166  65                  70                  75                  80
168 Phe Leu Ser Pro Leu Ala Ser Ser Ala Ala Ser Arg Ser Ser Ala Arg
169             85                  90                  95
171 Asp Asp Lys Leu Thr Ala Leu Leu Ala Gln Leu Asp Ser Leu Thr Arg
172             100                 105                 110
174 Glu Leu Asn Val Val Ser Gln Gln Leu Leu Asp Leu Arg Gln Gln Val
175             115                 120                 125
177 Ser Ala Leu Lys Ala Ser Ser Pro Pro Asn Ala Val
178      130             135             140
180 <210> SEQ ID NO: 6
181 <211> LENGTH: 132
182 <212> TYPE: PRT
183 <213> ORGANISM: Adenovirus
185 <400> SEQUENCE: 6
186 Met Ser Gly Phe Thr Glu Gly Asn Ala Val Ser Phe Glu Gly Gly Val
187   1                   5                   10                  15
189 Phe Ser Pro Tyr Leu Thr Thr Arg Leu Pro Ser Trp Ala Gly Val Arg
190             20                  25                  30
192 Gln Asn Val Val Gly Ser Asn Val Asp Gly Arg Pro Val Ala Pro Ala
193             35                  40                  45
195 Asn Ser Thr Thr Leu Thr Tyr Ala Thr Ile Gly Ser Ser Val Asp Thr
196      50                  55                  60
198 Ala Ala Ala Ala Ala Ala Ser Ala Ala Ala Ser Thr Ala Arg Gly Met
199  65                  70                  75                  80
201 Ala Ala Asp Phe Gly Leu Tyr Asn Gln Leu Ala Ala Ser Arg Leu Arg
202             85                  90                  95
204 Glu Glu Asp Ala Leu Ser Val Val Leu Thr Arg Leu Glu Glu Leu Ser
205             100                 105                 110
207 Gln Gln Leu Gln Asp Met Ser Ala Lys Met Ala Leu Leu Asn Pro Pro
208             115                 120                 125
210 Ala Asn Thr Ser
211      130
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Input Set : **A:\208859.txt**
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213 <210> SEQ ID NO: 7
214 <211> LENGTH: 133
215 <212> TYPE: PRT
216 <213> ORGANISM: Adenovirus
218 <400> SEQUENCE: 7
219 Met Ser Gly Ser Met Glu Gly Asn Ala Val Ser Phe Lys Gly Gly Val
220   1               5                  10                  15
222 Phe Ser Pro Tyr Leu Thr Thr Arg Leu Pro Ala Trp Ala Gly Val Arg
223             20                  25                  30
225 Gln Asn Val Met Gly Ser Asn Val Asp Gly Arg Pro Val Ala Pro Ala
226             35                  40                  45
228 Asn Ser Ala Thr Leu Thr Tyr Ala Thr Val Gly Ser Ser Val Asp Thr
229         50                  55                  60
231 Ala Ala Ala Ala Ala Ala Ser Ala Ala Ala Ser Thr Ala Arg Gly Met
232   65                  70                  75                  80
234 Ala Ala Asp Phe Gly Leu Tyr Asn Gln Leu Ala Ala Ser Arg Ser Leu
235             85                  90                  95
237 Arg Glu Glu Asp Ala Leu Ser Val Val Leu Thr Arg Met Glu Glu Leu
238             100                 105                 110
240 Ser Gln Gln Leu Gln Asp Leu Phe Ala Lys Val Ala Leu Leu Asn Pro
241             115                 120                 125
243 Pro Ala Asn Ala Ser
244       130
246 <210> SEQ ID NO: 8
247 <211> LENGTH: 130
248 <212> TYPE: PRT
249 <213> ORGANISM: Adenovirus
251 <220> FEATURE:
252 <221> NAME/KEY: misc_feature
253 <222> LOCATION: (2)..(5)
254 <223> OTHER INFORMATION: "Xaa" may be any amino acid
256 <220> FEATURE:
257 <221> NAME/KEY: misc_feature
258 <222> LOCATION: (7)..(7)
259 <223> OTHER INFORMATION: "Xaa" may be any amino acid
261 <220> FEATURE:
262 <221> NAME/KEY: misc_feature
263 <222> LOCATION: (9)..(9)
264 <223> OTHER INFORMATION: "Xaa" may be any amino acid
266 <220> FEATURE:
267 <221> NAME/KEY: misc_feature
268 <222> LOCATION: (11)..(11)
269 <223> OTHER INFORMATION: "Xaa" may be any amino acid
271 <220> FEATURE:
272 <221> NAME/KEY: misc_feature
273 <222> LOCATION: (21)..(21)
274 <223> OTHER INFORMATION: "Xaa" may be any amino acid
276 <220> FEATURE:
277 <221> NAME/KEY: misc_feature
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Input Set : **A:\208859.txt**
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278 <222> LOCATION: (30)..(30)
279 <223> OTHER INFORMATION: "Xaa" may be any amino acid
281 <220> FEATURE:
282 <221> NAME/KEY: misc_feature
283 <222> LOCATION: (34)..(34)
284 <223> OTHER INFORMATION: "Xaa" may be any amino acid
286 <220> FEATURE:
287 <221> NAME/KEY: misc_feature
288 <222> LOCATION: (45)..(45)
289 <223> OTHER INFORMATION: "Xaa" may be any amino acid
291 <220> FEATURE:
292 <221> NAME/KEY: misc_feature
293 <222> LOCATION: (54)..(57)
294 <223> OTHER INFORMATION: "Xaa" may be any amino acid
296 <220> FEATURE:
297 <221> NAME/KEY: misc_feature
298 <222> LOCATION: (69)..(69)
299 <223> OTHER INFORMATION: "Xaa" may be any amino acid
301 <220> FEATURE:
302 <221> NAME/KEY: misc_feature
303 <222> LOCATION: (74)..(76)
304 <223> OTHER INFORMATION: "Xaa" may be any amino acid
306 <220> FEATURE:
307 <221> NAME/KEY: misc_feature
308 <222> LOCATION: (79)..(83)
309 <223> OTHER INFORMATION: "Xaa" may be any amino acid
311 <220> FEATURE:
312 <221> NAME/KEY: misc_feature
313 <222> LOCATION: (86)..(86)
314 <223> OTHER INFORMATION: "Xaa" may be any amino acid
316 <220> FEATURE:
317 <221> NAME/KEY: misc_feature
318 <222> LOCATION: (88)..(90)
319 <223> OTHER INFORMATION: "Xaa" may be any amino acid
321 <220> FEATURE:
322 <221> NAME/KEY: misc_feature
323 <222> LOCATION: (92)..(95)
324 <223> OTHER INFORMATION: "Xaa" may be any amino acid
326 <220> FEATURE:
327 <221> NAME/KEY: misc_feature
328 <222> LOCATION: (98)..(98)
329 <223> OTHER INFORMATION: "Xaa" may be any amino acid
331 <220> FEATURE:
332 <221> NAME/KEY: misc_feature
333 <222> LOCATION: (100)..(101)
334 <223> OTHER INFORMATION: "Xaa" may be any amino acid
336 <220> FEATURE:
337 <221> NAME/KEY: misc_feature
338 <222> LOCATION: (105)..(105)
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VERIFICATION SUMMARY   DATE: 06/07/2001
PATENT APPLICATION: **US/09/780,224A**   TIME: 13:54:11

Input Set : **A:\208859.txt**
Output Set: **C:\CRF3\06072001\I780224A.raw**

```
L:377 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:380 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:383 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:386 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:389 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:395 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:401 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
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